

Form PTO-1449

Docket Number 511582003420

Application Number 10/087,190

INFORMATION DISCLOSURE CITATION
IN AN APPLICATION

(Use several sheets if necessary)

Applicant

Pia M. CHALLITA-EID et al.

Filing Date February 28, 2002

Group Art Unit 1642

Mailing Date December 16, 2004

U.S. PATENT DOCUMENTS

Examiner Initials	Ref. No.	Date	Document No.	Name	Class	Subclass	Filing Date If Appropriate
DB	1.	02/13/2003	2003/0032087	Challita-Eid et al.	11	11	
DB	2.	07/17/2001	6,262,333	Endege et al.	11	11	
DB	3.	05/27/2003	6,569,662	Tang et al.	11	11	

FOREIGN PATENT DOCUMENTS

Examiner Initials	Ref. No.	Date	Document No.	Country	Class	Subclass	Translation YES NO
DB	4.	07/08/1999	WO 99/33982	WIPO	11	11	
	5.	08/05/1999	WO 99/38972	WIPO	11	11	
	6.	12/16/1999	WO 99/64576	WIPO	11	11	
	7.	01/11/2001	WO 01/02568	WIPO	11	11	
	8.	04/05/2001	WO 01/22920	WIPO	11	11	
	9.	07/26/2001	WO 01/53312	WIPO	11	11	
	10.	09/13/2001	WO 01/66753	WIPO	11	11	
	11.	10/11/2001	WO 01/75067	WIPO	11	11	
	12.	10/24/2002	WO 02/083876	WIPO	11	11	
DB	13.	11/28/2002	WO 02/095009	WIPO	11	11	

OTHER DOCUMENTS

(including author, title, Date, Pertinent Pages, Etc.)

Examiner Initials	Ref. No.	Title
DB	14.	Attwood (2000). <i>Science</i> 290:471-473.
	15.	Burgess et al. (1990). <i>J. Cell Biol.</i> 111:2129-2138.
	16.	Carninci, P. et al. (1999). "High-Efficiency Full-Length cDNA Cloning" Chapter 2 <i>In cDNA Preparation and Characterization - Methods in Enzymology</i> , Weissman, S.M. ed. Academic Press, Inc. 303:19-44.
DB	17.	Carninci, P. et al. (2000). "Normalization and Subtraction of Cap-Trapper-Selected cDNAs to Prepare Full-Length cDNA Libraries for Rapid Discovery of New Genes," <i>Genome Research</i> 10:1617-1630.

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18.	DGENE Record Accession No. AAX98924 for WO 99/33982, two pages.		
19.	DGENE Record Accession No. AAX99020 for WO 99/33982, two pages.		
20.	EMBL: Accession No. AAM40043, created on October 21, 2001, European Search Report Reference XP 002297423, two pages		
21.	EMBL: Accession No. AYO28916, created on March 30, 2001, European Search Report Reference No. XP002297424, one page		
22.	EMBL: Accession No. AAZ80011 created on April 7, 2000, European Search Report Reference No. XP-002297421, one page		
23.	EMBL: Accession No. BE513408 created on August 8, 2000, European Search Report Reference No. XP-002297422, two pages		
24.	GenBank Accession No. BAB27765 created on April 3, 2004 located at < http://www.ncbi.nlm.nih.gov/entrez/viewer.fcgi?db=protein&val=12847934 > last visited on November 28, 2004, three pages.		
25.	GenBank Accession No. NP_115493 created on October 27, 2004 located at < http://www.ncbi.nlm.nih.gov/entrez/viewer.fcgi?db=protein&val=14149769 > last visited on November 28, 2004, two pages.		
26.	Hubert et al. (1999). <i>Proc. Natl. Acad. Sci. USA</i> 96(25):14523-14528.		
27.	Klein et al. (1997). <i>Nat. Med.</i> 3:402.		
28.	Lazar et al. (1988). <i>Mol. Cell. Biol.</i> 8:1247-1252.		
29.	Lerner, (1982). <i>Nature</i> 299:592-596.		
30.	Metzler et al. (1997). <i>Nature Structural Biol.</i> 4:527-531.		
31.	Mikayama et al. (1993). <i>PNAS</i> 90:10056-10060.		
32.	Ngo et al. (1994). <i>The Protein Folding Problem and Tertiary Structure Prediction</i> Birkhauser: Boston, MA pp. 433, 492-495.		
33.	Pinto et al. (1996). <i>Clin. Cancer Res.</i> 2(9):1445-1451.		
34.	Reiter et al. (1998). <i>Proc. Natl. Acad. Sci. USA</i> 95:1735.		
35.	Shibata, K. et al. (2000). "RIKEN Integrated Sequence Analysis (RISA) System - 384-Format Sequencing Pipeline with 384 Multicapillary Sequencer," <i>Genome Research</i> 10:1757-1771.		
36.	Skolnick et al. (2000). <i>Trends in Biotech.</i> 18(1):34-39.		
37.	Su et al. (1996). <i>Proc. Natl. Acad. Sci. USA</i> 93:7252.		
38.	Supplementary Partial Search Report mailed October 6, 2004, for European Patent Application No. 02723291.7, filed February 28, 2002, 7 pages.		
39.	The FANTOM Consortium and the RIKEN Genome Exploration Research Group Phase I & II Team (December 5, 2002). "Analysis of the Mouse Transcriptome Based on Functional Annotation of 60,770 Full-Length cDNAs," <i>Nature</i> 420:563-573.		
EXAMINER: <i>David B. Burt</i>		DATE CONSIDERED: <i>4/20/05</i>	
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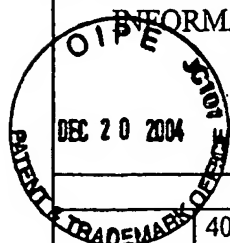
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40.

The RIKEN Genome Exploration Research Group Phase II Team and the FANTOM Consortium (February 8, 2001). "Functional Annotation of a Full-Length Mouse cDNA Collection," *Nature* 409:685-690.

41.

Tsubouchi, H. et al. (May 2002). "The Mnd1 Protein Forms a Complex with Hop2 to Promote Homologous Chromosome Pairing and Meiotic Double-Strand Break Repair," *Molecular and Cellular Biology* 22(9):3078-3088.

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